

THE RELATIONSHIP BETWEEN
HUMAN DIFFERENTIAL LEARNING/ADAPTIVE BEHAVIOR
AND AN INDICATOR OF COMPLEMENT REGULATION LYSIS

by

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THE RELATIONSHIP BETWEEN
HUMAN ERYTHROCYTE LIPIDOLYTISSOME RECEPTOR
AND AN INHIBITOR OF COMPLEMENT ACTIVATED LYSIS

By

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Human erythrocyte membrane extracts with receptor specificity
for the lipopolysaccharides of gram negative bacteria were found to
be rich in DR, a complement inhibitory substance originally isolated
and described from human red cell membranes. The possibility that
the DR-receptor and DR inhibitor might be the same macromolecule was
examined.

In this investigation it was demonstrated that although closely
associated, the two activities are biologically distinct and separable.
This conclusion was supported by the results of five experimental ap-
proaches. First, but distinct differences were observed in the elution
profiles of the two activities when crude extracts of erythrocyte mem-
branes were subjected to Sepharose 4B, G4M-Sepharose chromatography
and polyacrylamide thin gel electrophoresis. Complete separation of
the two activities was accomplished by a shift in the pH of the membrane
extraction medium. Further, differences between the two activities
based on their ability to spontaneously bind to sheep erythrocytes were

demonstrated. Finally, complement was not activated, indicating that the absence of sheep erythrocytes (E) instead of cells lacking LPS-receptor specificity but same density of IgG inhibitor activity.

Additional studies indicated that LPS bound to cell membranes can activate either the alternative or classical complement pathways and that IgG inhibitor associated with cell membranes can block LPS-induced complement lysis of red cells, despite the fact that complement activation has occurred.

INTRODUCTION

The lipopolysaccharides (LPS) of gram negative bacteria are among numerous antigens known to be capable of binding to the membranes of erythrocytes and other mammalian cells *in vitro* and under certain conditions, *in vivo* (1). Because of their unique ability to modulate the immune response in a wide variety of ways, they have emerged as a complex and fascinating class of macromolecules. Functionally, antibodies have been shown to have very different properties due to their chemical makeup and localization in the outer membrane of the bacterial cell envelope they have been shown to play a major role in the establishment of a selective permeability barrier (2,3) and in serving as receptors for certain bacteriophages (4).

Of interest to the researcher, however, is the fact that interactions of LPS with components of the immune system may lead to a single or combination of physiological responses. These include toxicity, antigenicity, immunogenicity, tolerance and activation of complement. Although much is known concerning the general nature of these responses, the mechanisms of the cell associated events responsible for their development in the presence of LPS are still not fully understood.

A great deal of information about the chemical structure of LPS, from a variety of organisms, has accumulated. Although it has been recently recognized that LPS isolated from a given organism is

heterogeneous (5). LPS of most gram-negative virulent organisms appear to share the same basic molecular composition. As illustrated in Figure 1, all consist of three regions. The first region, the O-specific polysaccharide section which is made up of repeating units of five to eight monosaccharides, carries the main serologic specificity for a given organism. Numerous serological groups differing in O-section specificity are now recognized and the polysaccharide accordingly shows wide inter- and intrageneric variations in composition (6). Of interest is the fact that natural antibodies to this region are found in most animal species but do not always appear to be protective, and in some cases a lethal gram-negative bacteremia develops despite high titers of O-specific antibody (1). Besides the presence of a short ester core which contains glucose [glc], galactose [gal], and 5-acetylglucosamine [GlcNAc], and an inner core of 1-glycosyl-2-ketodeoxyribose, phosphatidyl, ribitolamine and three molecules of 3-deoxyoctate (KDO). Unique to the antibodies of gram negatives, 3-deoxyoctate provides a linkage site to the third region, the third A section. Lipid A is basically composed of a phosphorylated glucosamine backbone to which are attached fatty acids and ethanolamine residues. The nature and distribution of the Lipid A fatty acids varies among bacterial groups with the inner core polysaccharide composition revealing some shift (7). The complete LPS containing O-specific antigen is designated somatic (8), and all mutants lacking O-specific side chains are referred to as rough or H-forms.

Many early studies concerned with the interaction of antibodies and biological systems were carried out using either whole bacteria



16

Abstract

Figure 1.

Protein structure of (A) of *Escherichia coli*. There are presently three proteins that function in protein synthesis. The ribosome is composed of two subunits, the large and small subunits. The large subunit contains the 23S and 23S ribosomal RNA molecules, which are the catalytic core of the ribosome. The small subunit contains the 16S and 16S ribosomal RNA molecules, which are the structural core of the ribosome. The ribosome is composed of two subunits, the large and small subunits. The large subunit contains the 23S and 23S ribosomal RNA molecules, which are the catalytic core of the ribosome. The small subunit contains the 16S and 16S ribosomal RNA molecules, which are the structural core of the ribosome. (A) map also to protein [4].

of interest to bacterial LPS preparations in the fluid phase. They are in spite of the fact that it had been shown that heated LPS (100°C for 80 minutes) could be taken into the surface of a number of cells including erythrocytes (8). The latter are often the model target cells for hemolytic assays. It has become increasingly clear that the biological consequences of either *in vivo* or *in vitro* encounter with LPS, whether cell bound or partially purified, is dependent upon its fixation to various target cells (3). For example, it was reported that patients suffering from endotoxin shock and sepsis due to gram negative bacteria had greatly reduced levels of blood platelets and that the platelets contained LPS. It has further been established that human platelets possess an endotoxin binding receptor, which when triggered by interaction with LPS, results in the release of a vasoactive amine, 5-hydroxytryptamine (5-HT) and the generation of clot-promoting activity (9). In animal studies, it was observed that guinea pigs injected intravenously with LPS also show a 50% decrease in blood platelets with a concomitant shortening of the clotting time (10). Additionally, it has been reported that there is a direct relationship between the susceptibility of different strains of mice to the lethal effects of *Salmonella enteritidis* and the affinity of their red cells for either heat-killed *Salmonella* or free LPS (12).

Little was known about the nature of the attachment of LPS to any cell until the iron citrate, strain 6, serotype isolated as extract from the membranes of human erythrocytes having a high affinity and specificity for the lipopolysaccharides of a variety of gram negative

SECRETIN [13-18]. This material, designated as an LPS-receptor, has now been purified to homogeneity and characterized. Spranger has reported that the LPS-receptor is a lipopolysaccharide, rich in 8-acetyl-saccharic acid (8ASA), galactose, fucose, sialic acid and carries about 475 proteins [14]. It appears to be a pentameric molecule with a molecular weight of about 250,000 daltons. The LPS-receptor functions by direct interaction with groups on the LPS molecule which provide an attachment site for tissue components [16]. Given evidence has accumulated suggesting that this attachment site is the lipid A moiety of LPS [17]. This high affinity of the LPS-receptor for endotoxin is quite remarkable because both macromolecules are highly negatively charged. The receptor, because of its high 8ASA content and LPS because of its phosphoric acid moieties.

Because the immunological specificity of LPS based on erythrocytes remains unchanged, Spranger has suggested that the lipid A of LPS binds to the specific receptor via clusters of hydrophobic amino acids which make up about 40% of the total peptide content of the receptor leaving the polysaccharide available for the reaction with antibodies [16]. A complete understanding of the orientation of LPS on tissues, bound either by specific receptors or by non-specific mechanisms, may come from studies involving the interaction of cell bound LPS with serum complement.

The auto-complementary effects of LPS have long been established. For some time, evidence seemed to suggest that the single most important factor in the development of a serious response to endotoxin was the direct interaction of the lipid A region with individual

systems including the complement system (14). Moreover, the mechanism, eliciting evidence seem to indicate complement as a mediator of a number of the deleterious responses to endotoxins in experimental animals and man (15). More importantly, recent evidence indicated that the ability of LPS to initiate a complement response is not confined to the Thiel-Auloy but appears to involve the polysaccharide core as well (16).

It has become increasingly clear that a major role of the complement system during an immune response is the amplification of ordered pleiotropies with a concomitant induction of an inflammatory response. This is accomplished by the sequential activation of the proteins which make up the complement system. The activation process can be divided into three major stages: a recognition stage, the generation of C3 cleavage enzymes and C3 activation stage, and a terminal or membrane attack stage (10-12). Under normal conditions, complement proteins exist in the serum as inactive precursors, and are activated by either of two pathways -- the classical or alternative. The components of these pathways and their reaction requirements are summarized in Table 1. The two share a number of similar characteristics but differ, notably, in the initiation and the sequence of reactions of the first two stages.

The capacity of endotoxins to activate the complement system by a mechanism which requires neither antibodies to LPS nor the participation of the early complement components was demonstrated more than twenty years ago (21). More recently, it was recognized that Thiel phase activation of complement by LPS is not restricted to the

TABLE I
Comparative Properties of the Classical
and Alternative Complement Pathways

	Classical	Alternative
Activating agents		
Immunochemical of human	IgG1-IgG3, IgA/IgM/IgE	IgM and IgG
cellular	IgM	IgM
serum pig	IgG2	IgG1
microbial	IgM	IgG2
Regulation	Factor D	Factor 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100
Activating site	Pro-Fragment	Pro-Fragment (Factor 1) or Pro-Fragment (Factor 2) or Pro-Fragment (Factor 3) or Pro-Fragment (Factor 4) or Pro-Fragment (Factor 5) or Pro-Fragment (Factor 6) or Pro-Fragment (Factor 7) or Pro-Fragment (Factor 8) or Pro-Fragment (Factor 9) or Pro-Fragment (Factor 10) or Pro-Fragment (Factor 11) or Pro-Fragment (Factor 12) or Pro-Fragment (Factor 13) or Pro-Fragment (Factor 14) or Pro-Fragment (Factor 15) or Pro-Fragment (Factor 16) or Pro-Fragment (Factor 17) or Pro-Fragment (Factor 18) or Pro-Fragment (Factor 19) or Pro-Fragment (Factor 20) or Pro-Fragment (Factor 21) or Pro-Fragment (Factor 22) or Pro-Fragment (Factor 23) or Pro-Fragment (Factor 24) or Pro-Fragment (Factor 25) or Pro-Fragment (Factor 26) or Pro-Fragment (Factor 27) or Pro-Fragment (Factor 28) or Pro-Fragment (Factor 29) or Pro-Fragment (Factor 30) or Pro-Fragment (Factor 31) or Pro-Fragment (Factor 32) or Pro-Fragment (Factor 33) or Pro-Fragment (Factor 34) or Pro-Fragment (Factor 35) or Pro-Fragment (Factor 36) or Pro-Fragment (Factor 37) or Pro-Fragment (Factor 38) or Pro-Fragment (Factor 39) or Pro-Fragment (Factor 40) or Pro-Fragment (Factor 41) or Pro-Fragment (Factor 42) or Pro-Fragment (Factor 43) or Pro-Fragment (Factor 44) or Pro-Fragment (Factor 45) or Pro-Fragment (Factor 46) or Pro-Fragment (Factor 47) or Pro-Fragment (Factor 48) or Pro-Fragment (Factor 49) or Pro-Fragment (Factor 50) or Pro-Fragment (Factor 51) or Pro-Fragment (Factor 52) or Pro-Fragment (Factor 53) or Pro-Fragment (Factor 54) or Pro-Fragment (Factor 55) or Pro-Fragment (Factor 56) or Pro-Fragment (Factor 57) or Pro-Fragment (Factor 58) or Pro-Fragment (Factor 59) or Pro-Fragment (Factor 60) or Pro-Fragment (Factor 61) or Pro-Fragment (Factor 62) or Pro-Fragment (Factor 63) or Pro-Fragment (Factor 64) or Pro-Fragment (Factor 65) or Pro-Fragment (Factor 66) or Pro-Fragment (Factor 67) or Pro-Fragment (Factor 68) or Pro-Fragment (Factor 69) or Pro-Fragment (Factor 70) or Pro-Fragment (Factor 71) or Pro-Fragment (Factor 72) or Pro-Fragment (Factor 73) or Pro-Fragment (Factor 74) or Pro-Fragment (Factor 75) or Pro-Fragment (Factor 76) or Pro-Fragment (Factor 77) or Pro-Fragment (Factor 78) or Pro-Fragment (Factor 79) or Pro-Fragment (Factor 80) or Pro-Fragment (Factor 81) or Pro-Fragment (Factor 82) or Pro-Fragment (Factor 83) or Pro-Fragment (Factor 84) or Pro-Fragment (Factor 85) or Pro-Fragment (Factor 86) or Pro-Fragment (Factor 87) or Pro-Fragment (Factor 88) or Pro-Fragment (Factor 89) or Pro-Fragment (Factor 90) or Pro-Fragment (Factor 91) or Pro-Fragment (Factor 92) or Pro-Fragment (Factor 93) or Pro-Fragment (Factor 94) or Pro-Fragment (Factor 95) or Pro-Fragment (Factor 96) or Pro-Fragment (Factor 97) or Pro-Fragment (Factor 98) or Pro-Fragment (Factor 99) or Pro-Fragment (Factor 100)
Factors required to generate C3 convertase	C1q C4 C2	Factor 1 Factor 2 Factor 3 Factor 4 Factor 5 Factor 6 Factor 7 Factor 8 Factor 9 Factor 10 Factor 11 Factor 12 Factor 13 Factor 14 Factor 15 Factor 16 Factor 17 Factor 18 Factor 19 Factor 20 Factor 21 Factor 22 Factor 23 Factor 24 Factor 25 Factor 26 Factor 27 Factor 28 Factor 29 Factor 30 Factor 31 Factor 32 Factor 33 Factor 34 Factor 35 Factor 36 Factor 37 Factor 38 Factor 39 Factor 40 Factor 41 Factor 42 Factor 43 Factor 44 Factor 45 Factor 46 Factor 47 Factor 48 Factor 49 Factor 50 Factor 51 Factor 52 Factor 53 Factor 54 Factor 55 Factor 56 Factor 57 Factor 58 Factor 59 Factor 60 Factor 61 Factor 62 Factor 63 Factor 64 Factor 65 Factor 66 Factor 67 Factor 68 Factor 69 Factor 70 Factor 71 Factor 72 Factor 73 Factor 74 Factor 75 Factor 76 Factor 77 Factor 78 Factor 79 Factor 80 Factor 81 Factor 82 Factor 83 Factor 84 Factor 85 Factor 86 Factor 87 Factor 88 Factor 89 Factor 90 Factor 91 Factor 92 Factor 93 Factor 94 Factor 95 Factor 96 Factor 97 Factor 98 Factor 99 Factor 100
Final serum requirement	Optimal	Concentrated
Final serum requirements	C4 and C2	C3

^aC1 is a trypsin-like complex of C1q, C1r and C1s. Classical pathway activation is initiated when C1q binds to immunoglobulin.

Classical pathway activation of LPS have been demonstrated to activate the classical pathway in either the presence or absence of specific antibodies (3,24-25,27).

Normally, during classical pathway activation, the recognition stage is initiated by the fixation of the first component of the complement system, C1, to IgG or IgM immunoconjugates of specific immune complexes. This fixation and activation of C1 generates a C1 esterase whose major substrates are the fourth and second components of the complement system.

Endotoxins in the fluid phase have been shown to initiate the activation of the complement system in vitro as described above in the presence of antibody or IgM antibodies directed against the α -polysaccharide antigen (28). In spite of the initiation of certain gram negatives, however, is the capacity of the lipid A moiety of the molecule to fix C1 directly resulting in a antibody independent initiation of the classical complement cascade (3,28).

Alternative pathway activation, on the other hand, is triggered by direct activation of an Initiator Factor (IF) by antibodies, or any one of a number of naturally occurring polysaccharides, and aggregates of IgG or IgE (29). It has been established that alternative pathway activation of the complement system involves the core polysaccharide region of the LPS molecule (30). Additionally, it has been shown that the length of the O-specific polysaccharide antigen and differences in the carbohydrate content may also play a role in this mechanism of activation (31).

In short, two of the classical pathway activation routes, C3 convertase is generated when C3 esterase cleaves C4 and C2 in the presence of negative particles (the cell bound C3aC2 complex). This convertase cleaves the third component of complement (C3) into two fragments, C3a and C3b. The latter fragment is unstable and has a highly relative hydrophobic binding site. As a result of binding to target molecules at sites adjacent to the molecules bound C3 convertase, a new enzyme C5 convertase, whose major substrate becomes the fifth component of complement, is generated. In addition, C3b undergoes secondary changes giving rise to an immune-adhesiveness site capable of binding to a variety of effector cells of the immune system which bear specific C3b receptors (32). Additionally, C3b, either bound hydrophobically to a cell surface or free in solution, can produce further splitting of C3 via the C3 feedback cycle of the alternative pathway described below. It is this amplification of the generation of C3b and its deposition via immune adhesiveness onto the surface of specific target cells (and in some cases basement membranes), where the major function of the complement system is realized.

The mechanism of the alternative pathway generation of the C3 convertase is a bit more complicated in that there is a direct requirement for preformed C3b, the source of which is still not fully understood. This C3b is connected with factor B plus properdin and factor D from the alternative pathway convertase "Factor BCD" capable of splitting C3 into C3a and C3b (33). This split was active C3 giving rise to further C3b and the cycle is again repeated. This reaction

is regulation dependent and while classical pathway activation is inhibited by high concentrations of calcium [34], factor proper converts neutral conditions these critical requirements make serum diluted with either ethylenediamine tetraacetic acid (EDTA) or ethylenedipyl tetraacetic acid (EGTA) useful reagents for distinguishing between the two pathways [35]. The former, being an effective chelator of both calcium and magnesium blocks the activation of the two pathways, while the latter, a less effective chelator of magnesium, preferentially blocks the classical pathway.

The first and final stage of complement activation is the same for both pathways and is initiated with the cleavage of C3 into two fragments, the real C3b by the C3 convertase. The larger C3a fragment then reacts sequentially with C4 and C2 to form either a cell bound or fluid phase trimolecular complex C567. The cell bound complex has the capability to bind C5 and C6. If the cell to which this C5-C6 complex is associated is sensitive to complement mediated cytotoxicity, lysis ensues [36].

Fluid phase C567, C5a, and C5b are potent opolylectins and chemotaxis [37,38]. Once leukocytes such as polymorphonuclear leukocytes (PMN) and macrophages have migrated to the site of complement activation, phagocytosis is initiated. As previously stated, the phagocytic process is enhanced by the fixation of complement components, especially C3b, onto the surface of particulate antigens or target tissues with protein adherence, thus facilitating ingestion [39]. Release of Haptocytolipid complexes, either as a direct consequence of ingestion or expulsion of an unphagocytized target into the surrounding tissue, results in the generation of additional chemotactic factors (40,41). Thus

model of complement regulation, the early stage in the development of an antibody response.

Several clinical studies resulting from extended application of the activation of the complement system is severely inhibited by the presence of two human serum inhibitors, C1 esterase inhibitor (C1I) and complement activating factor (CAF) also known as C2b inactivator (C2bI). Besides there are other humoral or cellular inhibitory factors which is to be defined.

Many of the studies cited here employed fresh plasma LPS for *in vitro* assays. Neter was the first to demonstrate that LPS coated onto the surface of sheep erythrocytes were sensitive to lysis by an antibody dependent classical complement pathway mechanism (44). Phillips and Bengtson also studying these erythrocytes treated with LPS subjected by the heat-plate-plate procedure (45) confirmed Neter's observation showing the need for the presence of a naturally occurring γ_2 globulin for classical activation with a splitting observation of the early components C1-C3 (46).

The *in vivo* consequences of the intravenous administration of endotoxins to normal and complement deficient animals have been studied. Both complement pathways are activated with the classical pathway being involved for the development of many of the pathophysiological responses such as the thrombocytopenia observed in many experimental endotoxin infections (11). Some investigators have suggested that classical pathway activation due to an indirectly mediated function of LPS to various cells such as platelets and erythrocytes is more effective in

activation (some) systems with membrane-associated inhibitor release of a variety of cytotoxic agents for phagocytosis (27,44). Alternative pathway activation has not been shown to inhibit the release of membrane damaging factors (48). In vivo systems are difficult to evaluate because of the large number of parameters which must be considered. Therefore, much study is needed before a complete understanding of the reactions involved can be obtained.

Regardless of the nature of the system (in vitro or in vivo) or the state of the endotoxin (free or cell bound), the interaction of LPS with the complement system in the presence of specific antibody is extremely efficient, giving full response with serum concentrations of antibody and C1-C3 (44). Resistance of some negative bacteria would then appear to be related to the facts and rate of free endotoxin released as a direct consequence of cell death due to phagocytosis and lysis or uptake of this free endotoxin with other cell systems in the presence of serum factors.

As previously stated, erythrocytes and also other cells have been shown to have membrane receptors which bind endotoxin. The molecular role of these receptors is still not clear, however, fixation of LPS to the cell does appear to be a precondition for the triggering of many mechanisms of the immune system to LPS (48).

Recently, in the late 1960s, isolated and described extracts from the membranes of human erythrocytes capable of inhibiting the hemolytic activity of complement when adsorbed onto erythrocytes were used as target cells (46). Extracts isolated at two different localities

cytolytic potential of C5b6b is being attributed for hemolysis of sheep erythrocytes. Material prepared at a molar stoichiometry of 6:1 was shown to be capable of lysis in sheep erythrocytes and protecting them from osmotic lysis, and has been designated as inhibitor.

An extract prepared under the same conditions but at a lower molarity was incapable of lysis in sheep erythrocytes but was capable of accelerating the decay of the complement component intermediates $C4C2$ to $C4C4$. This material was designated I_{II} for decay acceleration factor (II).

Preliminary studies indicate that the I_{II} inhibitor is a large molecule with a molecular weight greater than 250,000. It appears to contain at least two sugar moieties, glucose and galactose and it is about 45 proteins by weight (52).

Existing data support that the I_{II} inhibitor probably acts at the $C3$ conversion step. This was suggested by the finding that I_{II} coated sheep erythrocytes in the intermediate state $C3C4$ consumed less $C3$ than untreated controls and that the inhibitory effects of I_{II} coated once activated $C3$ became fixed to the cell bound $C3$ convertase ($C3, C3$).

Attempts to more fully define the biochemical and biological properties of these intermediates have been hampered by the inability to obtain them in a highly purified state.

Erythrocytes of different species differ in their susceptibility as target cells to immune hemolysis, with sheep and chicken erythrocytes being far more sensitive than human and guinea pig erythrocytes. Differences within the same species have also been observed (53). For

1980) it has been reported that patients suffering from paroxysmal nocturnal hemoglobinuria (PNH) have at least two populations of erythrocytes tested on their susceptibility to immune hemolysis (IH), with at least one subpopulation exhibiting extreme sensitivity to attack by an antibody-independent complement mediated mechanism. It has been demonstrated that extracts from the erythrocyte stroma of these patients have reduced levels of Df inhibitor activity.¹ Additionally, evidence has been accumulated which suggests that there is a parallel between the presence of DAF on the surfaces of the erythrocytes of certain patients and resistance to the cytolytic effects of complement (C5).

A comparison of the isolation schemes for purifying the Df inhibitor and the UPE-receptor revealed a marked similarity between the two both activities are confined to that fraction of the membrane extractable by a heptanol-water mixture at 4°C and at an ionic strength of 2 M or below. All of the activity of either material appeared to be localized only in the aqueous layer of the heptanol extractant supernatant (an exchange chromatography of extracts containing either the UPE-receptor or Df inhibitor activities indicates that both are eluted under identical conditions). These observations would suggest that the two activities may be similar or even identical.

The principle objective of this investigation was to determine if the Df inhibitor and UPE-receptor are either the same or closely

¹Data supported by personal experiments

related. Although it has been observed that human erythrocytes are highly resistant to LPS mediated immune hemolysis, the reason for the refractoriness of these treated erythrocytes has not been defined. Sheep erythrocytes, as previously stated, are normally sensitive to immune lysis but may be rendered resistant by treatment with diluted extracts of the partially purified D₁ inhibitor. Therefore, the second objective of this investigation was to explore the biological consequences of the D₁ inhibitor to the interaction of LPS treated erythrocytes and serum complement, in an attempt to clearly establish a biological role for the D₁ inhibitor.

MATERIALS AND METHODS

Erythrocytes. Sub-arterial human blood (Group B, Rh positive) containing erythritol-phosphate-dehydrogenase as an enzyme marker was obtained from the Dr. H. H. Regional Blood Center (Gainesville, FL). Whole (group blood was taken by venipuncture from subjects hospitalized at the Animal Research Laboratory of the U. S. H. H. Miller Health Center (Gainesville, FL). One volume of blood was mixed with an equal volume of sterile heparinized Alsever's solution (36) and the blood was stored at 4°C for up to one month.

Expansion of erythrocyte stroma. Human and sheep erythrocyte stroma were prepared by the method of Springer et al. (36). Erythrocytes from whole blood were pelleted at 4°C by centrifugation for ten minutes at 500g and the plasma and buffy coat were discarded. The packed cells were washed three times with phosphate buffered saline (0.15M sodium chloride plus 0.005M potassium phosphate) at pH 7.4 and lysed in 50 volumes of distilled water at 4°C. In the initial studies, the pH was adjusted to 5.3 with acetic acid and phenol was added to a final volume of 3.0%. The stroma were allowed to settle overnight at 4°C and the supernatant fluid was removed. Ten volumes of cold distilled water were added, and the pH was readjusted to 5.3. The stroma were sedimented either by settling or centrifugation and the entire procedure was repeated six times with the addition of phenol

Figure 2

A schematic representation of the calibration and verification procedures that were applied to the recordings at the LPA recorder of Springer and the 16 channels of Hoffman.

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reduced temperature (10%
depression in melting behavior)
increased water (1.5) sorption
in air, 4°C, 24 h (20)
activity in various media

100

the 1990s, the number of people in the United States who are 65 years of age or older is projected to increase from 20 million to 35 million, and the number of people 75 years of age or older is projected to increase from 10 million to 15 million (U.S. Census Bureau, 1996).

100

[illegible]Schubert, Bernhard. 1994. *Lehrbuch der Zoologie*. 10th ed. Berlin: Springer.

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2001 volume, starting at 10 pages.
Many thanks to J. B. and con-
tributors. 2001-2002
Revisions 2002 volume, 2003 edition,
at 34, 10 to only volume

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after early seeded water change. Following the final wash the strainer was collected by centrifugation, weighed and stored at -80°C until use.

Isolation and purification of the LPS-receptor. The LPS-receptor was prepared as described in Figure 2 using a modification of the procedure of Springer et al. [11]. A 100 species streptomycin suspension was homogenized in a starting blender and extracted overnight with two volumes of n-butyl alcohol at 4°C for 16 hours at pH 8.2. Four phases, resulting after centrifugation at 3000g for 30 minutes, appeared, namely, 1)top, aqueous and white. The aqueous phase was re-extracted twice with n-butanol, once for 30 minutes and again overnight and then thoroughly dialyzed against several changes of 0.05 M Tris-Cl buffer [pH 7.0]. An aqueous butanol extract low in LPS-receptor activity but high in 18 inhibitor activity was obtained by shifting the pH of the butanol extraction from 8.2 to 5.2.

The dialyzed active crude butanol extract was centrifuged at 150,000 g for 1-2 hours in a Spinco Model L5 preparative ultracentrifuge. Three phases resulted from the high speed centrifugation. Contrary to Springer's findings, the top aqueous layer possessed the highest LPS-receptor. After extensive dialysis, the aqueous top layer was applied to a 50 x 1.5 cm Sepharose 6B column (Pharmacia Fine Chemicals, Piscataway, N.J.). The Sepharose column was washed with a 0.05 M Tris-Cl buffer at pH 7.0. Three millionth fractions were collected and assayed for both LPS-receptor and 18 inhibitor activities. The active performing fractions were pooled, concentrated tenfold by dialysis against 30% polyethylene-glycol in 0.05 M Tris-Cl buffer [pH 7.0] and applied to a 20.0 x 2.5 cm

DEAE-Sepharose 4B (Pharmacia Fine Chemicals, Piscataway, NJ) column. After extensive washing of the column with the starting buffer, a linear sodium chloride gradient was initiated with 150 ml of 0.05 M pH 7.5 Tris-HCl buffer and 150 ml of 0.25 M, pH 7.5 Tris-HCl buffer. Three milliliter fractions were collected and assayed for the two activities.

Other extractions. Sometimes LPH-receptor or binding activity could not be localized in the aqueous bottom phase after shift in the pH during extraction. The intermediate lipid phase was then further extracted with equal volumes of ether (Phillipsburgh, NJ, Lotis, 40). Equal volumes of the lipid phase, suspended in 0.05 M Tris-HCl buffer at a pH of 7.5 (3.0) and ether were vigorously mixed in a separatory funnel for 5-15 minutes. After phase separation the ether was removed from the aqueous and spray intermediate lipids by diluting against phosphate buffered saline (pH 7.4) and from the organic phase by evaporation using a stream of nitrogen at 4°C. Following evaporation, the residue remaining from the organic phase was resuspended in the original volume with PBS and all phases were tested for LPH-receptor and IB inhibitor activities.

Polyacrylamide gel electrophoresis (PAGE). Disc polyacrylamide electrophoresis was performed using a modification of the method of Hunter [37]. Extracts were applied to 2.5% acrylamide gels and were electrophoresed in a non-reducing Tris-glycine buffer system, pH 8.8, for 18 minutes at 4°C. The gels were stained with 0.02% Coomassie blue containing 12.5% trichloroacetic acid. For some studies, duplicate gels were run. One gel was stained as above with the remaining gel being stored for analysis for LPH-receptor and IB inhibitor activities.

Lyophilized LPS extracted by the Baum (28) and Mesrobian (29) procedures were purchased from Sigma Laboratories (Detroit, MI). Bacterial (100°C for 2 hours) and washed LPS stock solutions (0.5 mg/ml PBS at pH 7.4) were stored at -80°C until used.

Antisera Appropriate dilutions of Salmonella group 1 O:antiserum obtained from Wellcome Biological Laboratories (Woburn, MA) were made in 0.01 M PBS, pH 7. Hemagglutination (40) titers of the sera ranged from 64 to 256.

For hemolytic assays, rabbit red antibodies to sheep erythrocyte stromata were obtained from Ceder Laboratories (Miami, FL). Stock solutions at a dilution of 1:1000 in PBS (pH 7.4) were maintained until use at -80°C.

Incubation of erythrocytes with LPS Freshly acquired erythrocytes from a group A, 15 positive adult were obtained and used in the coating and coating inhibition assays which were carried out as described by Springer et al. (14). For speed and economy, screening assays were assessed using a microplate hemagglutination system. Briefly, the procedure consisted of mixing equal volumes of either human or sheep erythrocytes at 2×10^8 cells/ml and dilutions of LPS for 45 minutes with shaking at 30°C. After extensive washing, the optimal amount of LPS which afforded maximal hemagglutination by subsequently added antiserum was determined. This dilution, defined as the optimal coating dose, was used in all subsequent hemagglutination-inhibition assays.

(25-membrane/100000 cells) LPS-receptor activity was determined by measuring the ability of a material to inhibit LPS fixation to yeast cells. The procedure in the coating inhibition assay differed from that in the coating test in that dilutions of LPS binding material were added to equal volumes of an optimal coating dose of LPS and incubated with shaking for 30 minutes at 37°C. Erythrocytes were added, and 18 filters described as previously described. In each assay, a control consisting of LPS and erythrocytes but no LPS-receptor material followed by the subsequent addition of antiserum was included.

Isotonic buffer selected utilized in qualitative assays The basic diluent for most hemolytic assays was the isotonic gelatin serum buffer (ISB) described by Iselt and Seger [18] which contained 0.005% K_2CaCl_2 , 0.005% $\text{K}_2\text{H}_2\text{P}_2\text{O}_7$, and 0.1% gelatin at pH 7.5. In some cases, gelatin serum without CaCl_2 or $\text{K}_2\text{H}_2\text{P}_2\text{O}_7$, containing enough Tris-hydroxyacetate acetate (THAA, pH 7.4) to bring the final concentration to either 0.01 M or 0.04 M, was employed. These buffers were described as 0.01 M (THA-SB) and 0.04 M (THA-SB) respectively. In order to achieve maximal sensitivity, hemolytic assays involving individual complement components, and 18 inhibitor assays were performed using a low ionic strength gelatin-serum prepared by mixing equal volumes of 0.02 M Tris-hydroxyacetate buffer containing twice the standard amount of CaCl_2 and $\text{K}_2\text{H}_2\text{P}_2\text{O}_7$ (ISB).

Enriched sheep erythrocytes [18] Fresh erythrocytes at a concentration of 10^8 per ml in 0.01 M THA-SB were mixed with an equal volume of actively to sheep serum at a final dilution of 1:100 in

stop flow buffer. The C11(1)Pm was incubated with shaking for 30 minutes at 37°C, and then at 0°C for either 30 minutes or overnight. The cells were washed twice and standardized to the desired concentration before use.

Complement (C5). Fresh frozen guinea pig complement was obtained from Bell Trans Laboratories (Napier, AB). The serum was diluted to dry ice and was stored at -20°C until use. In some studies, aliquots of C5 serum were absorbed three times at 0°C with either untreated or LPS treated sheep or human erythrocytes before use.

Complement Components. Guinea pig C1 and C2 were prepared by methods described by Nilsson et al. [30] and Fiedler and Davin [35,31] respectively. Individual purified functionally pure guinea pig complement components C3, C5, C6, C7, C8 and C9 were purchased from Cohn Laboratories (Peters, PA).

Guinea pig complement intermediates. For C3 inhibition assays and complement concentration studies, sheep erythrocytes in the intermediate state C3i, C5i7, and C5i7i2 were prepared by the methods of Nilsson and Sage [32].

Identification of form of C5i7i2. The identity of the generation of C5i7i2 was determined by the Test procedure described by Nilsson et al. [33].

C3 inhibition experiments. Cross human erythrocyte stroma extracts, high in C3 inhibitor activity, were isolated by a procedure described by Hoffmann [34]. The method is outlined in Figure 2. Briefly, the essential differences to Hoffmann's preparation of isolated

Highly purified quantity (45% to 55% inhibitor activity) and extracts high in LPS-receptor activity as defined by Jeffries are: (1) streams were prepared at pH of 8.0 - 7.5 without the addition of bleach; (2) crude washed streams were adjusted to equal values of 0.005 M potassium phosphate buffer at a pH of 7.5 and extracted with chloroform at a final concentration of 20% for 15 minutes; and, (3) after the first extraction the aqueous chloroform phase was adjusted to an ionic strength of 0.15 by the addition of 0.5 M NaCl. Reduced extraction of the adjusted material was repeated until a light phase could no longer be separated. For further purification, concentrated material, active or in activity, was subjected to gel filtration and HPLC chromatography.

Treatment of sheep erythrocytes with artificially modified lipopolysaccharide. Lipopolysaccharide isolated and extracted sheep red blood erythrocytes were treated with Dn inhibitor material by the procedure described by Jeffries (50). Equal volumes of erythrocytes at 10^8 cells/ml to 0.005 ml extracts of Dn inhibitor labeled 1.18 to 0.005 were mixed at 4°C. The mixture was transferred to a 30°C water bath, incubated 30 minutes with shaking and was pelleted at 5000g for 10 minutes at 4°C. The pellet cells were washed two times, and standardized to the desired concentration in the appropriate buffer.

Dn inhibitor activity. Dn inhibitor activity was assessed using the OACPE inactivation assay described by Jeffries (50). Sheep erythrocytes in the intermediate state (OACPE) at 10^8 /ml were mixed with an equal volume of Dn inhibitor material of target in 0.005. The mixture was incubated at 30°C for 15 minutes with constant shaking,

After 30 min, three volumes of guinea pig complement diluted 1:25 in 0.2 M KSCN-BS were added. The tubes were then incubated for 60 min more at 37°C with shaking. At the end of the incubation period, 10 volumes of ice-cold PBS was added to each reaction mixture. The cells were pelleted at 500g for 10 minutes at 4°C and the optical densities of the supernatant fluids were determined at a wave length of 411 nm.

Reagents. Stock solutions of reaction ethyleneimide-labeled bovine HMB, Fisher Scientific Co., Fair Lawn, NJ and ethyleneimide-labeled [carboxymethyl ether] α^2 , β mannanside acid (BZA, Sigma Chemical Co., St. Louis, MO) were prepared as described by Fine et al. (28). The stock solutions were stored at 4°C and diluted to a final concentration of 200 μ M before use. Reaction BZA was prepared as described by Fine et al. (28).

Complement Consumption. The ability of erythrocytes treated with LPS and/or D inhibitor to consume complement was determined in reaction mixtures containing 0.1 ml of the treated cells (1×10^8 cells) or LPS and 0.5 ml of normal or absorbed guinea pig serum challenged with either BSA or BZA. The mixtures were incubated with shaking 60 minutes at 37°C. Following the incubation period, the cells were pelleted at 500g at 4°C for 10 minutes. The supernatant fluids were reconstituted with negative control cells and were analyzed for residual whole complement activity using a modification of the procedure as outlined by Gehai and Fayer (29).

Erythrocyte coating by LPS and its inhibition Repeated titrations

of Salmonella typhimurium LPS at concentrations ranging from 0.001 μg to 10.0 $\mu\text{g}/\text{ml}$ as determined with polyvalent and homologous Salmonella antisera, employing erythrocytes at 2.5×10^8 cells/ml were carried out. The results of a representative experiment employing polyvalent antisera are given in table II. It can be seen that heating the LPS enhanced the erythrocyte coating capacity to a remarkable extent. Additionally, a maximum titer resulted when erythrocytes were exposed to at least 2.50 $\mu\text{g}/\text{ml}$ of heated LPS. Therefore, an optimal coating unit (u) of heated LPS (defined as the reciprocal of the greatest dilution of LPS producing complete hemagglutination by either polyvalent or homologous antisera) was taken as 0.10 $\mu\text{g}/\text{ml}$. Results employing homologous antisera to heated LPS yielded lower optimal coating doses of 0.10 $\mu\text{g}/\text{ml}$ and 1.04 $\mu\text{g}/\text{ml}$, depending on the age of the antisera. These results were identical for human and sheep erythrocytes. LPS extracted by the Vesigal procedure resulted in an optimal coating dose of 0.10 $\mu\text{g}/\text{ml}$ as also determined with homologous antisera.

LPS receptor activity, as evaluated in these studies, was based on the ability of a given erythrocyte preparation to inhibit the complete fixation of an optimal coating unit of LPS onto either sheep or human

TABLE 1

Determination of optimal LPS concentration
used for coating human erythrocytes

LPS concn	Erycys ^a	
	untreated	coated ^b
10.0	80	80
7.5	110	80
5.0	60	80
2.5	40	80
0.125	60	100
0.063	60	100
0.031	40	100
0.016	20	80
0.008	10	80

^aThe percentage of the addition of anti-LPS serum affecting maximal hemagglutination

^b100% additions of LPS (1.0 mg/ml PBS) were tested (20°C for three hours)

cytotoxicity. Table III, below, summarizes the results obtained with several crude aqueous ethanol preparations of erythrocyte stroma. These data indicate that the range of LPS-receptor concentrations or activities needed to yield optimal inhibition of LPS-erythrocyte coating varied with the source, concentration and condition (pH/heat) of the erythrocyte stroma extraction procedure.

(B) Inhibitor activity of crude ethanol extracts of human and sheep erythrocyte stroma. OAGT_2 Inactivation by a crude erythrocyte stroma extract, prepared by the technique described in the section on materials and methods is shown in Figure 5. This procedure was used to determine the inhibitory efficacy of most extracts. Color controls for the presence of hemoglobin in the higher concentrations of crude preparations were necessary. The reciprocal of the dilution of a crude ethanol preparation yielding greater than 50% inhibition of the lysis of OAGT_2 by C-DEA are also shown in Table III. These results clearly indicate that LPS-receptor and its inhibitor activities are contained in significant amounts in crude ethanol stroma extracts obtained by either Springer's or Raffelson's procedures. Of interest also, is that the potency of the two activities varied to the same extent.

A comparison of the physicochemical properties of the its inhibitor and LPS-receptor from erythrocyte stroma extracts. The above data suggests that the LPS-receptor and its inhibitor are either identical or closely related molecules, therefore, additional evidence to resolve this issue was sought. Human and sheep erythrocyte stroma were subjected to modifications of Springer's purification procedure as outlined

Table III

Comparison of LPS-reactor and II activity
obtained in several Crude Extracts of *Hydrophaga* species

Crude Extracts (Species)	Titers	
	LPS-reactor ^a	II activity ^b
Roses cells extracted at pH 8.2 (Hoffmann)		
Crude aqueous butanol phase	40	40
Crude aqueous phase		
high speed ^c top layer	160	120
high speed x-rayp interface	80	40
Roses cells extracted at pH 7.5 (Hoffmann)		
Crude aqueous butanol phase	160	120
Roses cells extracted at pH 7.2 (Sprague)		
Crude aqueous butanol phase	0	0
Crude aqueous ether phase	0	0
Crude ether interface	0	0
Crude ether aqueous phase	0	0

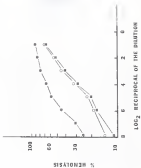
^aThe smallest amount (diffusion) giving complete inhibition of LPS reacting

^bThe reciprocal of the greatest dilution giving 50% inhibition of OCHT
[unit]

^cHigh speed extracts were obtained by the centrifugation of the crude
aqueous butanol phase, 10,000g for 2 hours

Figure 3.

Isolation of the mutants of *lacT* by D-6275 is the presence of various concentrations of above grade protein resulting from the lactose repression of *lac* operon. *lac* operon mutants appeared according to the presence of lactose. The spot clusters due to activity of the *lac* operon (lac⁺ mutant) showed that *lac* operon cluster than the lactation associated with the *lac* protein appeared after the high speed centrifugation (40,000 rpm/15 min) of the crude extracted samples. The cluster appeared above the higher activity of lactation obtained after the high speed centrifugation of the extracted samples.



In Figure 2, Figure 4 is a gel filtration elution profile on sepharose 4B, of the crude high speed top layer obtained from the crude latent large myelocyte stromal preparation. Fractions were monitored at 220 and 280 nm and were assayed for LPS-receptor and 24 inhibitor activities as previously described. Two peaks were observed with both activities eluting in the peak following the void volume. Close examination of the sepharose 4B profile indicates that there is a slight displacement of the 24 inhibitor activity to the left of the LPS-receptor activity. This would suggest that perhaps the two activities may be different.

Further attempts to separate and purify the two activities were accomplished using ion exchange chromatography. The sepharose 4B active peaks were pooled, dialyzed against the starting Tris-HCl buffer, and applied to a BSA-Sepharose column. Fractionation was accomplished with a linear NaCl gradient. A typical demonstration of the partially purified material(s) is shown in Figure 5. LPS-receptor and 24 inhibitor activities eluted in a relatively narrow peak at about 0.1M NaCl, again with the 24 inhibitor slightly preceding the LPS-receptor activity.

The recovery of the LPS-receptor and 24 inhibitor activities following sepharose 4B and BSA-Sepharose chromatography is presented in Table 17. It should be noted that gel filtration on sepharose 4B yielded only about one-half increase in the purity of both activities with recovery of only 50% of the LPS-receptor activity and 40% of the 24 inhibitor activity. BSA-Sepharose was shown to result in as much as a 10-fold increase in activity, but resulted in a recovery of 80% of the specific LPS-receptor activity but only one third or 33% of the

TABLE 11

Recovery of UTP-enzyme and its residual activity from deposits of human erythrocyte membranes following separation of lipid-deposits. Percentage

Preparation of lipid-deposits	[wt] (g/g ^a)	Concentration (mg/ml)	Activated (mg)	Total Enzyme activity (mg)	Specific activity (mg)	Total activity (mg)	Residual activity (mg)	Total activity (mg)
Separation of lipid								
Dried high lipid (100%)	475.2	320	200	1,150	1,440	1,440	1,440	1,440
Separation of recovery of UTP								
100%	10.5	10	20	200	200	200	200	200
50%-lipid-deposits								
100%	10	100	80	40	240	440	440	440

^aThe residual amount (in mg) of the lipid-deposits of UTP coating

^bThe percentage of the residual activity of UTP coating

^cCalculation

^dCalculation of dividing the value for total activity by the value for total concentration

^eCalculation of dividing the value for total activity after treatment, by the value for total concentration

^fCalculation of dividing the value for each of the residual activities by the value for the total activity

Figure 4

2. (a) Effluents resulting from the (b) (c) extracted human activities
 were samples prepared by the method of (d) (e) at all the
 extracted material were applied to a sequence of columns (f) (g)
 (h) (i) and chromatographed at 25°C with 0.1M HCl buffer, pH
 1.0. (j) (k) (l) (m) (n) (o) (p) (q) (r) (s) (t) (u) (v) (w) (x) (y) (z)
 (aa) (ab) (ac) (ad) (ae) (af) (ag) (ah) (ai) (aj) (ak) (al) (am) (an) (ao) (ap) (aq) (ar) (as) (at) (au) (av) (aw) (ax) (ay) (az) (ba) (bb) (bc) (bd) (be) (bf) (bg) (bh) (bi) (bj) (bk) (bl) (bm) (bn) (bo) (bp) (bq) (br) (bs) (bt) (bu) (bv) (bw) (bx) (by) (bz) (ca) (cb) (cc) (cd) (ce) (cf) (cg) (ch) (ci) (cj) (ck) (cl) (cm) (cn) (co) (cp) (cq) (cr) (cs) (ct) (cu) (cv) (cw) (cx) (cy) (cz) (da) (db) (dc) (dd) (de) (df) (dg) (dh) (di) (dj) (dk) (dl) (dm) (dn) (do) (dp) (dq) (dr) (ds) (dt) (du) (dv) (dw) (dx) (dy) (dz) (ea) (eb) (ec) (ed) (ee) (ef) (eg) (eh) (ei) (ej) (ek) (el) (em) (en) (eo) (ep) (eq) (er) (es) (et) (eu) (ev) (ew) (ex) (ey) (ez) (fa) (fb) (fc) (fd) (fe) (ff) (fg) (fh) (fi) (fj) (fk) (fl) (fm) (fn) (fo) (fp) (fq) (fr) (fs) (ft) (fu) (fv) (fw) (fx) (fy) (fz) (ga) (gb) (gc) (gd) (ge) (gf) (gg) (gh) (gi) (gj) (gk) (gl) (gm) (gn) (go) (gp) (gq) (gr) (gs) (gt) (gu) (gv) (gw) (gx) (gy) (gz) (ha) (hb) (hc) (hd) (he) (hf) (hg) (hh) (hi) (hj) (hk) (hl) (hm) (hn) (ho) (hp) (hq) (hr) (hs) (ht) (hu) (hv) (hw) (hx) (hy) (hz) (ia) (ib) (ic) (id) (ie) (if) (ig) (ih) (ii) (ij) (ik) (il) (im) (in) (io) (ip) (iq) (ir) (is) (it) (iu) (iv) (iw) (ix) (iy) (iz) (ja) (jb) (jc) (jd) (je) (jf) (jg) (jh) (ji) (jj) (jk) (jl) (jm) (jn) (jo) (jp) (jq) (jr) (js) (jt) (ju) (jv) (jw) (jx) (jy) (jz) (ka) (kb) (kc) (kd) (ke) (kf) (kg) (kh) (ki) (kj) (kk) (kl) (km) (kn) (ko) (kp) (kq) (kr) (ks) (kt) (ku) (kv) (kw) (kx) (ky) (kz) (la) (lb) (lc) (ld) (le) (lf) (lg) (lh) (li) (lj) (lk) (ll) (lm) (ln) (lo) (lp) (lq) (lr) (ls) (lt) (lu) (lv) (lw) (lx) (ly) (lz) (ma) (mb) (mc) (md) (me) (mf) (mg) (mh) (mi) (mj) (mk) (ml) (mm) (mn) (mo) (mp) (mq) (mr) (ms) (mt) (mu) (mv) (mw) (mx) (my) (mz) (na) (nb) (nc) (nd) (ne) (nf) (ng) (nh) (ni) (nj) (nk) (nl) (nm) (nn) (no) (np) (nq) (nr) (ns) (nt) (nu) (nv) (nw) (nx) (ny) (nz) (oa) (ob) (oc) (od) (oe) (of) (og) (oh) (oi) (oj) (ok) (ol) (om) (on) (oo) (op) (oq) (or) (os) (ot) (ou) (ov) (ow) (ox) (oy) (oz) (pa) (pb) (pc) (pd) (pe) (pf) (pg) (ph) (pi) (pj) (pk) (pl) (pm) (pn) (po) (pp) (pq) (pr) (ps) (pt) (pu) (pv) (pw) (px) (py) (pz) (qa) (qb) (qc) (qd) (qe) (qf) (qg) (qh) (qi) (qj) (qk) (ql) (qm) (qn) (qo) (qp) (qq) (qr) (qs) (qt) (qu) (qv) (qw) (qx) (qy) (qz) (ra) (rb) (rc) (rd) (re) (rf) (rg) (rh) (ri) (rj) (rk) (rl) (rm) (rn) (ro) (rp) (rq) (rr) (rs) (rt) (ru) (rv) (rw) (rx) (ry) (rz) (sa) (sb) (sc) (sd) (se) (sf) (sg) (sh) (si) (sj) (sk) (sl) (sm) (sn) (so) (sp) (sq) (sr) (ss) (st) (su) (sv) (sw) (sx) (sy) (sz) (ta) (tb) (tc) (td) (te) (tf) (tg) (th) (ti) (tj) (tk) (tl) (tm) (tn) (to) (tp) (tq) (tr) (ts) (tt) (tu) (tv) (tw) (tx) (ty) (tz) (ua) (ub) (uc) (ud) (ue) (uf) (ug) (uh) (ui) (uj) (uk) (ul) (um) (un) (uo) (up) (uq) (ur) (us) (ut) (uu) (uv) (uw) (ux) (uy) (uz) (va) (vb) (vc) (vd) (ve) (vf) (vg) (vh) (vi) (vj) (vk) (vl) (vm) (vn) (vo) (vp) (vq) (vr) (vs) (vt) (vu) (vv) (vw) (vx) (vy) (vz) (wa) (wb) (wc) (wd) (we) (wf) (wg) (wh) (wi) (wj) (wk) (wl) (wm) (wn) (wo) (wp) (wq) (wr) (ws) (wt) (wu) (wv) (ww) (wx) (wy) (wz) (xa) (xb) (xc) (xd) (xe) (xf) (xg) (xh) (xi) (xj) (xk) (xl) (xm) (xn) (xo) (xp) (xq) (xr) (xs) (xt) (xu) (xv) (xw) (xx) (xy) (xz) (ya) (yb) (yc) (yd) (ye) (yf) (yg) (yh) (yi) (yj) (yk) (yl) (ym) (yn) (yo) (yp) (yq) (yr) (ys) (yt) (yu) (yv) (yw) (yx) (yz) (za) (zb) (zc) (zd) (ze) (zf) (zg) (zh) (zi) (zj) (zk) (zl) (zm) (zn) (zo) (zp) (zq) (zr) (zs) (zt) (zu) (zv) (zw) (zx) (zy) (zz)

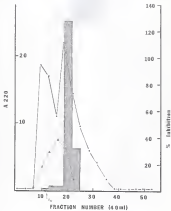
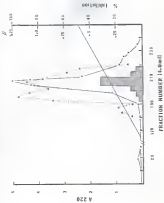


Figure 5

A 5000-Scanline A-25 chromatophore profile of the sediment at active station two from Ulmunda estuary prepared by computer of Figure 4.3 of text material. With an optical density at 200 m of 10.8 was applied to a 20 x 20 m section. A 5 hour median filtering process of 0.25 x 0.25 m filters, at 10 m applied and a 10 m filtering was reduced. Section divided at the station at 200 m and shown by the closed circles. The closed triangles indicate the 5 hour median of the station capable of filtering both at deep section and the closed bars represent the divisions, having 1/20-median activity.



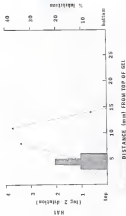
comparing IL inhibitor activity in comparison of the IL inhibitor and OAD-leptanin elution profiles (Figures 4 and 5) and the data in Table II, revealing a 50% loss in the total area but with a 10-fold increase in activity, indicate good purification of the two activities. These data do, however, suggest further differences in the two activities by the differences in their respective recoveries following both reverse IL and OAD-leptanin chromatography.

The homogeneity of the active OAD-leptanin fractions was checked by electrophoresis on 7.5% polyacrylamide gels at pH 8.8. As shown in Figure 6, 4 - 12 protein bands were observed on the crude extracts with a sharp band and a diffused staining area localized at the top of the gel when partially purified OAD-leptanin fractions were electrophoresed. Analysis of duplicate gels indicated that most of the LPS-receptor activity was localized in an area about 4 cm into the gels with the IL inhibitor activity spread over a fairly large area at the top of the gel with a peak of activity at about 11.5 cm. (Figure 6)

IL inhibitor and LPS-receptor activities of sheep erythrocyte ghosts. The aqueous phase of butanol extracted sheep erythrocyte ghosts, prepared by Springer's extraction procedure at pH 8.2 and at pH 5.2, had neither detectable LPS receptor nor IL inhibitor activities. LPS-receptor activity but no IL inhibitor was, however, observed in an ether soluble fraction of the n-butanol/1-triol phase as was shown in Table III.

Figure 1.

Correlation between the distribution of LPR-receptor and its inhibitory activities from extracts of human erythrocytic stromata when subjected to polyvinylpyrrolidone and dextran chromatography. Extracts were applied to Sephadex T-10 gels and every elution-fractionated in a non-reducing 1% dioxane buffer. All 8 hr eluates at 20% after dextran-chromatography, were not tested for profiles with erythrocytic stroma and the stroma was not taken separately. Sequential eluates were eluted and analyzed for its inhibitor and LPR-receptor activities. The closed brackets indicate the elution position of the inhibitor and the shaded bars represent the elution profile of the LPR-receptor activity.



A significant difference in the two preparations was evident when C-49, previously observed that a shift in the pH from 8.2 to 5.2 of the acetone-ethanol extract of the crude erythrocyte stroma resulted in a preparative separation of little or no detectable LPS-receptor activity, but high to 18 inhibitor activity. The elution profile of the 18 inhibitor activity following ion exchange chromatography on DEAE-Sephadex of the specially purified material, as shown in Figure 7, was similar to that observed for material extracted at pH 8.2. LPS-receptor activity assays were not carried out on the remaining ethanol phases because these materials had been discarded before the impact of these observations were realized.

A shift in the pH to 5.2 of a crude ethanol extract of erythrocyte stroma obtained at a pH of 8.2 affected neither the LPS-receptor nor 18 inhibitor activities. This suggested the possibility that the LPS-receptor activity of the material extracted at a pH of 8.2 was not destroyed but was probably reprecipitated into another phase. Stroma extractions were carried out at pH 5.2 employing smaller volumes of erythrocyte stroma, using Levine's procedure, in an attempt to fractionate the LPS-receptor activity. As shown in Table 8, some 18 inhibitor activity was observed in the crude aqueous ethanol phase when 10-6 ml of packed stroma were extracted. There was no detectable LPS-receptor in this ethanol layer. The third phase mixed with 10 volumes of PEI was further subjected to an ether extraction resulting in four layers: a pellet, an aqueous layer, light etherphase, and an organic layer. All of the LPS-receptor activity was recovered in the largest ether layer

TABLE 7

Recovery of LPS-mono- and disaccharide activity of *Bacteroides* strains isolated at pH 5.5 after 48 h incubation at 37°C

Preparation	Quantity	Total Concentration (M_{100°)	Total Activity U/g ^a	Total Activity U/g ^b	Specific Activity U/g ^c
Respect. reduced	10.0 ml	18.0	0	72.0	0
Respect. reduced	10.0 ml	50.0	0	0	0
Lyophil	1.0 g	50.0			
Aqueous extract	0.5	0.10	0	100	0.0001
Lipid extract	0.5	0.0	0	100	0.0001
Respect. extract	10.0	100.0	100		0.0001

^aThe medium (medium) during medium incubation at 37°C activity.

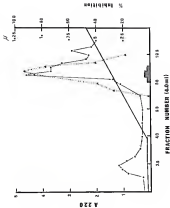
^bRecovery of the respect. activity during 48 h incubation at 37°C.

^cNot determined, due to high levels of particulate matter.

distributed by dividing the value for total activity by the value for total concentration.

Figure 2.

A, 300; B, pseudo A–D; chromatographic profiles of the solvent extracted from *Cratichneumon stramineus* prepared at pH 5.5 for the analysis of serotonin. 40.0 at of the initial material with an injection quantity of 100 nm of 1.0 were applied to a 10 x 0.5 cm column. A–D in fractions were eluted with a linear gradient elution profile of 0.05–0.75% TMS buffer at 0.5 mL/min. Optical densities of the fractions of 100 nm are shown by the closed circles. The closed triangles indicate the elution position of the enantiomeric isomers of octadecyl triethylsilyl ether (OC/TES) and the closed bars represent the fraction during LC–MS/MS activity.



with an average sedimentation coefficient of 100S and 100S supernatants. A comparison of the purification tables (IV and VI) and these extracts obtained at pH 8.2 and pH 5.5 indicates that, as was observed at pH 8.2, extracts obtained at pH 5.5 and subjected to DEAE-Sepharose chromatography resulted in a substantially greater loss of total mass as estimated by the absorbance at 225 m μ and the yield of 18 inhibitor activity.

Treatment of Serflinger's crude labeled extracts with sheep erythrocytes (226). To further establish that the two activities are distinctly different, equal volumes of sheep erythrocytes at either 10^8 or 10^9 cells/ml were mixed with equal volumes of a labeled high speed too layer extract prepared according to the procedure of Serflinger. This extract has an initial LPS-receptor titer of 128 and a 18g γ inhibitor titer greater than 80. Control tubes consisting of equal volumes of buffer and the extracts were also prepared. All tubes were mixed at 30°C for 30 minutes with shaking. The cells were pelleted by centrifugation and the supernatant fluids along with the buffer controls were diluted and assayed for LPS-receptor and 18 inhibitor activities. As can be seen in Figure 2, the 18 inhibitor activity was reduced substantially when extracts were treated with 10^8 cells/ml. In contrast, the LPS-receptor activity remained constant when the cell treated and buffer control supernatant fluids were compared.

The biological consequences of the 18 inhibitor and LPS-receptor on the erythrocyte membrane. The data in the previous section indicated two essential agents. First, the membrane of the human erythrocyte,

Figure 4

Treatment of cortical laminar neurons in sheep erythrocyte vesicles prepared by the method of Springer. Either 10^6 sheep erythrocytes or half the volume mixed with an equal volume of the untreated material. Following a 30 minute incubation period at 37°C , the cell suspension was pelleted and all supernatant fluids were assayed for UFB-receptor and IR substrate activities. The hatched bars represent the sheep erythrocyte treated material, and the open bars the buffer treated material.

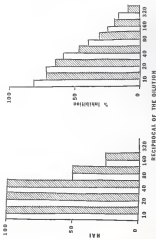


TABLE 21

Activity of 15 inhibitors and LPS-receptor activation
from *Cryptosporidium parvum* (isolated at pH 5.2) by the presence
of bovine and subjected to SDS-PAGE electrophoresis

Inhibitor	Conc.	Dilution	LPS-act ^a	LPS-act ^b	LPS-act ^c	LPS-act ^d	LPS-act ^e	LPS-act ^f	LPS-act ^g	LPS-act ^h	LPS-act ⁱ	LPS-act ^j	LPS-act ^k	LPS-act ^l	LPS-act ^m	LPS-act ⁿ	LPS-act ^o	LPS-act ^p	LPS-act ^q	LPS-act ^r	LPS-act ^s	LPS-act ^t	LPS-act ^u	LPS-act ^v	LPS-act ^w	LPS-act ^x	LPS-act ^y	LPS-act ^z	LPS-act ^{aa}	LPS-act ^{ab}	LPS-act ^{ac}	LPS-act ^{ad}	LPS-act ^{ae}	LPS-act ^{af}	LPS-act ^{ag}	LPS-act ^{ah}	LPS-act ^{ai}	LPS-act ^{aj}	LPS-act ^{ak}	LPS-act ^{al}	LPS-act ^{am}	LPS-act ^{an}	LPS-act ^{ao}	LPS-act ^{ap}	LPS-act ^{aq}	LPS-act ^{ar}	LPS-act ^{as}	LPS-act ^{at}	LPS-act ^{au}	LPS-act ^{av}	LPS-act ^{aw}	LPS-act ^{ax}	LPS-act ^{ay}	LPS-act ^{az}	LPS-act ^{ba}	LPS-act ^{bb}	LPS-act ^{bc}	LPS-act ^{bd}	LPS-act ^{be}	LPS-act ^{bf}	LPS-act ^{bg}	LPS-act ^{bh}	LPS-act ^{bi}	LPS-act ^{bj}	LPS-act ^{bk}	LPS-act ^{bl}	LPS-act ^{bm}	LPS-act ^{bn}	LPS-act ^{bo}	LPS-act ^{bp}	LPS-act ^{bq}	LPS-act ^{br}	LPS-act ^{bs}	LPS-act ^{bt}	LPS-act ^{bu}	LPS-act ^{bv}	LPS-act ^{bw}	LPS-act ^{bx}	LPS-act ^{by}	LPS-act ^{bz}	LPS-act ^{ca}	LPS-act ^{cb}	LPS-act ^{cc}	LPS-act ^{cd}	LPS-act ^{ce}	LPS-act ^{cf}	LPS-act ^{cg}	LPS-act ^{ch}	LPS-act ^{ci}	LPS-act ^{cj}	LPS-act ^{ck}	LPS-act ^{cl}	LPS-act ^{cm}	LPS-act ^{cn}	LPS-act ^{co}	LPS-act ^{cp}	LPS-act ^{cq}	LPS-act ^{cr}	LPS-act ^{cs}	LPS-act ^{ct}	LPS-act ^{cu}	LPS-act ^{cv}	LPS-act ^{cw}	LPS-act ^{cx}	LPS-act ^{cy}	LPS-act ^{cz}	LPS-act ^{da}	LPS-act ^{db}	LPS-act ^{dc}	LPS-act ^{dd}	LPS-act ^{de}	LPS-act ^{df}	LPS-act ^{dg}	LPS-act ^{dh}	LPS-act ^{di}	LPS-act ^{dj}	LPS-act ^{dk}	LPS-act ^{dl}	LPS-act ^{dm}	LPS-act ^{dn}	LPS-act ^{do}	LPS-act ^{dp}	LPS-act ^{dq}	LPS-act ^{dr}	LPS-act ^{ds}	LPS-act ^{dt}	LPS-act ^{du}	LPS-act ^{dv}	LPS-act ^{dw}	LPS-act ^{dx}	LPS-act ^{dy}	LPS-act ^{dz}	LPS-act ^{ea}	LPS-act ^{eb}	LPS-act ^{ec}	LPS-act ^{ed}	LPS-act ^{ee}	LPS-act ^{ef}	LPS-act ^{eg}	LPS-act ^{eh}	LPS-act ^{ei}	LPS-act ^{ej}	LPS-act ^{ek}	LPS-act ^{el}	LPS-act ^{em}	LPS-act ^{en}	LPS-act ^{eo}	LPS-act ^{ep}	LPS-act ^{eq}	LPS-act ^{er}	LPS-act ^{es}	LPS-act ^{et}	LPS-act ^{eu}	LPS-act ^{ev}	LPS-act ^{ew}	LPS-act ^{ex}	LPS-act ^{ey}	LPS-act ^{ez}	LPS-act ^{fa}	LPS-act ^{fb}	LPS-act ^{fc}	LPS-act ^{fd}	LPS-act ^{fe}	LPS-act ^{ff}	LPS-act ^{fg}	LPS-act ^{fh}	LPS-act ^{fi}	LPS-act ^{fj}	LPS-act ^{fk}	LPS-act ^{fl}	LPS-act ^{fm}	LPS-act ^{fn}	LPS-act ^{fo}	LPS-act ^{fp}	LPS-act ^{fq}	LPS-act ^{fr}	LPS-act ^{fs}	LPS-act ^{ft}	LPS-act ^{fu}	LPS-act ^{fv}	LPS-act ^{fw}	LPS-act ^{fx}	LPS-act ^{fy}	LPS-act ^{fz}	LPS-act ^{ga}	LPS-act ^{gb}	LPS-act ^{gc}	LPS-act ^{gd}	LPS-act ^{ge}	LPS-act ^{gf}	LPS-act ^{gg}	LPS-act ^{gh}	LPS-act ^{gi}	LPS-act ^{gj}	LPS-act ^{gk}	LPS-act ^{gl}	LPS-act ^{gm}	LPS-act ^{gn}	LPS-act ^{go}	LPS-act ^{gp}	LPS-act ^{gq}	LPS-act ^{gr}	LPS-act ^{gs}	LPS-act ^{gt}	LPS-act ^{gu}	LPS-act ^{gv}	LPS-act ^{gw}	LPS-act ^{gx}	LPS-act ^{gy}	LPS-act ^{gz}	LPS-act ^{ha}	LPS-act ^{hb}	LPS-act ^{hc}	LPS-act ^{hd}	LPS-act ^{he}	LPS-act ^{hf}	LPS-act ^{hg}	LPS-act ^{hh}	LPS-act ^{hi}	LPS-act ^{hj}	LPS-act ^{hk}	LPS-act ^{hl}	LPS-act ^{hm}	LPS-act ^{hn}	LPS-act ^{ho}	LPS-act ^{hp}	LPS-act ^{hq}	LPS-act ^{hr}	LPS-act ^{hs}	LPS-act ^{ht}	LPS-act ^{hu}	LPS-act ^{hv}	LPS-act ^{hw}	LPS-act ^{hx}	LPS-act ^{hy}	LPS-act ^{hz}	LPS-act ^{ia}	LPS-act ^{ib}	LPS-act ^{ic}	LPS-act ^{id}	LPS-act ^{ie}	LPS-act ^{if}	LPS-act ^{ig}	LPS-act ^{ih}	LPS-act ⁱⁱ	LPS-act ^{ij}	LPS-act ^{ik}	LPS-act ^{il}	LPS-act ^{im}	LPS-act ⁱⁿ	LPS-act ^{io}	LPS-act ^{ip}	LPS-act ^{iq}	LPS-act ^{ir}	LPS-act ^{is}	LPS-act ^{it}	LPS-act ^{iu}	LPS-act ^{iv}	LPS-act ^{iw}	LPS-act ^{ix}	LPS-act ^{iy}	LPS-act ^{iz}	LPS-act ^{ja}	LPS-act ^{jb}	LPS-act ^{jc}	LPS-act ^{jd}	LPS-act ^{je}	LPS-act ^{jf}	LPS-act ^{jj}	LPS-act ^{jk}	LPS-act ^{jl}	LPS-act ^{jm}	LPS-act ^{jn}	LPS-act ^{jo}	LPS-act ^{jp}	LPS-act ^{jq}	LPS-act ^{jr}	LPS-act ^{js}	LPS-act ^{jt}	LPS-act ^{ju}	LPS-act ^{jv}	LPS-act ^{jw}	LPS-act ^{jx}	LPS-act ^{ky}	LPS-act ^{kz}	LPS-act ^{la}	LPS-act ^{lb}	LPS-act ^{lc}	LPS-act ^{ld}	LPS-act ^{le}	LPS-act ^{lf}	LPS-act ^{lg}	LPS-act ^{lh}	LPS-act ^{li}	LPS-act ^{lj}	LPS-act ^{lk}	LPS-act ^{ll}	LPS-act ^{lm}	LPS-act ^{ln}	LPS-act ^{lo}	LPS-act ^{lp}	LPS-act ^{lq}	LPS-act ^{lr}	LPS-act ^{ls}	LPS-act ^{lt}	LPS-act ^{lu}	LPS-act ^{lv}	LPS-act ^{lw}	LPS-act ^{lx}	LPS-act ^{ly}	LPS-act ^{lz}	LPS-act ^{ma}	LPS-act ^{mb}	LPS-act ^{mc}	LPS-act ^{md}	LPS-act ^{me}	LPS-act ^{mf}	LPS-act ^{mg}	LPS-act ^{mh}	LPS-act ^{mi}	LPS-act ^{mj}	LPS-act ^{mk}	LPS-act ^{ml}	LPS-act ^{mm}	LPS-act ^{mn}	LPS-act ^{mo}	LPS-act ^{mp}	LPS-act ^{mq}	LPS-act ^{mr}	LPS-act ^{ms}	LPS-act ^{mt}	LPS-act ^{mu}	LPS-act ^{mv}	LPS-act ^{mw}	LPS-act ^{mx}	LPS-act ^{my}	LPS-act ^{mz}	LPS-act ^{na}	LPS-act ^{nb}	LPS-act ^{nc}	LPS-act nd	LPS-act ^{ne}	LPS-act ^{nf}	LPS-act ^{ng}	LPS-act ^{nh}	LPS-act ⁿⁱ	LPS-act ^{nj}	LPS-act ^{nk}	LPS-act ^{nl}	LPS-act ^{nm}	LPS-act ⁿⁿ	LPS-act ^{no}	LPS-act ^{np}	LPS-act ^{nq}	LPS-act ^{nr}	LPS-act ^{ns}	LPS-act ^{nt}	LPS-act ^{nu}	LPS-act ^{nv}	LPS-act ^{nw}	LPS-act ^{nx}	LPS-act ^{ny}	LPS-act ^{nz}	LPS-act ^{oa}	LPS-act ^{ob}	LPS-act ^{oc}	LPS-act ^{od}	LPS-act ^{oe}	LPS-act ^{of}	LPS-act ^{og}	LPS-act ^{oh}	LPS-act ^{oi}	LPS-act ^{oj}	LPS-act ^{ok}	LPS-act ^{ol}	LPS-act ^{om}	LPS-act ^{on}	LPS-act ^{oo}	LPS-act ^{op}	LPS-act ^{oq}	LPS-act ^{or}	LPS-act ^{os}	LPS-act ^{ot}	LPS-act ^{ou}	LPS-act ^{ov}	LPS-act ^{ow}	LPS-act ^{ox}	LPS-act ^{oy}	LPS-act ^{oz}	LPS-act ^{pa}	LPS-act ^{pb}	LPS-act ^{pc}	LPS-act ^{pd}	LPS-act ^{pe}	LPS-act ^{pf}	LPS-act ^{pg}	LPS-act ^{ph}	LPS-act ^{pi}	LPS-act ^{pj}	LPS-act ^{pk}	LPS-act ^{pl}	LPS-act ^{pm}	LPS-act ^{pn}	LPS-act ^{po}	LPS-act ^{pp}	LPS-act ^{pq}	LPS-act ^{pr}	LPS-act ^{ps}	LPS-act ^{pt}	LPS-act ^{pu}	LPS-act ^{pv}	LPS-act ^{pw}	LPS-act ^{px}	LPS-act ^{py}	LPS-act ^{pz}	LPS-act ^{qa}	LPS-act ^{qb}	LPS-act ^{qc}	LPS-act ^{qd}	LPS-act ^{qe}	LPS-act ^{qf}	LPS-act ^{qg}	LPS-act ^{qh}	LPS-act ^{qi}	LPS-act ^{qj}	LPS-act ^{qk}	LPS-act ^{ql}	LPS-act ^{qm}	LPS-act ^{qn}	LPS-act ^{qo}	LPS-act ^{qp}	LPS-act ^{qq}	LPS-act ^{qr}	LPS-act ^{qs}	LPS-act ^{qt}	LPS-act ^{qu}	LPS-act ^{qv}	LPS-act ^{qw}	LPS-act ^{qx}	LPS-act ^{qy}	LPS-act ^{qz}	LPS-act ^{ra}	LPS-act ^{rb}	LPS-act ^{rc}	LPS-act rd	LPS-act ^{re}	LPS-act ^{rf}	LPS-act ^{rg}	LPS-act ^{rh}	LPS-act ^{ri}	LPS-act ^{rj}	LPS-act ^{rk}	LPS-act ^{rl}	LPS-act ^{rm}	LPS-act ^{rn}	LPS-act ^{ro}	LPS-act ^{rp}	LPS-act ^{rq}	LPS-act ^{rr}	LPS-act ^{rs}	LPS-act ^{rt}	LPS-act ^{ru}	LPS-act ^{rv}	LPS-act ^{rw}	LPS-act ^{rx}	LPS-act ^{ry}	LPS-act ^{rz}	LPS-act ^{sa}	LPS-act ^{sb}	LPS-act ^{sc}	LPS-act ^{sd}	LPS-act ^{se}	LPS-act ^{sf}	LPS-act ^{sg}	LPS-act ^{sh}	LPS-act ^{si}	LPS-act ^{sj}	LPS-act ^{sk}	LPS-act ^{sl}	LPS-act sm	LPS-act ^{sn}	LPS-act ^{so}	LPS-act ^{sp}	LPS-act ^{sq}	LPS-act ^{sr}	LPS-act ^{ss}	LPS-act st	LPS-act ^{su}	LPS-act ^{sv}	LPS-act ^{sw}	LPS-act ^{sx}	LPS-act ^{sy}	LPS-act ^{sz}	LPS-act ^{ta}	LPS-act ^{tb}	LPS-act ^{tc}	LPS-act ^{td}	LPS-act ^{te}	LPS-act ^{tf}	LPS-act ^{tg}	LPS-act th	LPS-act ^{ti}	LPS-act ^{tj}	LPS-act ^{tk}	LPS-act ^{tl}	LPS-act tm	LPS-act ^{tn}	LPS-act ^{to}	LPS-act ^{tp}	LPS-act ^{tq}	LPS-act ^{tr}	LPS-act ^{ts}	LPS-act ^{tt}	LPS-act ^{tu}	LPS-act ^{tv}	LPS-act ^{tw}	LPS-act ^{tx}	LPS-act ^{ty}	LPS-act ^{tz}	LPS-act ^{ua}	LPS-act ^{ub}	LPS-act ^{uc}	LPS-act ^{ud}	LPS-act ^{ue}	LPS-act ^{uf}	LPS-act ^{ug}	LPS-act ^{uh}	LPS-act ^{ui}	LPS-act ^{uj}	LPS-act ^{uk}	LPS-act ^{ul}	LPS-act ^{um}	LPS-act ^{un}	LPS-act ^{uo}	LPS-act ^{up}	LPS-act ^{uq}	LPS-act ^{ur}	LPS-act ^{us}	LPS-act ^{ut}	LPS-act ^{uu}	LPS-act ^{uv}	LPS-act ^{uw}	LPS-act ^{ux}	LPS-act ^{uy}	LPS-act ^{uz}	LPS-act ^{va}	LPS-act ^{vb}	LPS-act ^{vc}	LPS-act ^{vd}	LPS-act ^{ve}	LPS-act ^{vf}	LPS-act ^{vg}	LPS-act ^{vh}	LPS-act ^{vi}	LPS-act ^{vj}	LPS-act ^{vk}	LPS-act ^{vl}	LPS-act ^{vm}	LPS-act ^{vn}	LPS-act ^{vo}	LPS-act ^{vp}	LPS-act ^{vq}	LPS-act ^{vr}	LPS-act ^{vs}	LPS-act ^{vt}	LPS-act ^{vu}	LPS-act ^{vv}	LPS-act ^{vw}	LPS-act ^{vx}	LPS-act ^{vy}	LPS-act ^{vz}	LPS-act ^{wa}	LPS-act ^{wb}	LPS-act ^{wc}	LPS-act ^{wd}	LPS-act ^{we}	LPS-act ^{wf}	LPS-act ^{wg}	LPS-act ^{wh}	LPS-act ^{wi}	LPS-act ^{wj}	LPS-act ^{wk}	LPS-act ^{wl}	LPS-act ^{wm}	LPS-act ^{wn}	LPS-act ^{wo}	LPS-act ^{wp}	LPS-act ^{wq}	LPS-act ^{wr}	LPS-act ^{ws}	LPS-act ^{wt}	LPS-act ^{wu}	LPS-act ^{wv}	LPS-act ^{ww}	LPS-act ^{wx}	LPS-act ^{wy}	LPS-act ^{wz}	LPS-act ^{xa}	LPS-act ^{xb}	LPS-act ^{xc}	LPS-act ^{xd}	LPS-act ^{xe}	LPS-act ^{xf}	LPS-act ^{xg}	LPS-act ^{xh}	LPS-act ^{xi}	LPS-act ^{xj}	LPS-act ^{xk}	LPS-act ^{xl}	LPS-act ^{xm}	LPS-act ^{xn}	LPS-act ^{xo}	LPS-act ^{xp}	LPS-act ^{xq}	LPS-act ^{xr}	LPS-act ^{xs}	LPS-act ^{xt}	LPS-act ^{xu}	LPS-act ^{xv}	LPS-act ^{xw}	LPS-act ^{xx}	LPS-act ^{xy}	LPS-act ^{xz}	LPS-act ^{ya}	LPS-act ^{yb}	LPS-act ^{yc}	LPS-act ^{yd}	LPS-act ^{ye}	LPS-act ^{yf}	LPS-act ^{yg}	LPS-act ^{yh}	LPS-act ^{yi}	LPS-act ^{yj}	LPS-act ^{yk}	LPS-act ^{yl}	LPS-act ^{ym}	LPS-act ^{yn}	LPS-act ^{yo}	LPS-act ^{yp}	LPS-act ^{yq}	LPS-act ^{yr}	LPS-act ^{ys}	LPS-act ^{yt}	LPS-act ^{yu}	LPS-act ^{yv}	LPS-act ^{yw}	LPS-act ^{yx}	LPS-act ^{yy}	LPS-act ^{yz}	LPS-act ^{za}	LPS-act ^{zb}	LPS-act ^{zc}	LPS-act ^{zd}	LPS-act ^{ze}	LPS-act ^{zf}	LPS-act ^{zg}	LPS-act ^{zh}	LPS-act ^{zi}	LPS-act ^{zj}	LPS-act ^{zk}	LPS-act ^{zl}	LPS-act ^{zm}	LPS-act ^{zn}	LPS-act ^{zo}	LPS-act ^{zp}	LPS-act ^{zq}	LPS-act ^{zr}	LPS-act ^{zs}	LPS-act ^{zt}	LPS-act ^{zu}	LPS-act ^{zv}	LPS-act ^{zw}	LPS-act ^{zx}	LPS-act ^{zy}	LPS-act ^{zz}

which (1) bind, resistant to complement mediated lysis, possess a, however, a distinctly different molecules with biologically varying properties. Endopolysaccharides with a high affinity for blood sugar saccharides which are potent activators of the complement system, and a class of molecules shown to be potent inactivators of complement. Sheep cells which are normally extremely sensitive to immune lysis have been shown to be devoid of the D₉ inhibitor but possess molecules with an affinity for LPS which are confined to the fluid moiety of the membrane.

As shown in Figure 3, the interaction of free LPS (extracted by both the Birnboim and Wisthaler procedures) with guinea pig serum which had been absorbed with stromal coated with LPS (S-LPS), resulted in a substantial consumption of complement. Additionally, it can be seen that LPS (Birboim) appeared to be a much more efficient activator of the alternative complement pathway compared to LPS extracted by the Wisthaler procedure.

LPS (extracted by both procedures) coated onto the surfaces of sheep erythrocytes showed a similar profile (Table III), except erythrocytes coated with LPS extracted by the Wisthaler procedure were far more efficient activators of complement in the absence of natural antibodies to LPS. It was of interest, therefore, to determine if LPS on the surface of sheep erythrocytes, in the presence of the D₉ inhibitor and S-LPS absorbed guinea pig serum, would alter the complement consumption profile of S-LPS. To explore this possibility, erythrocytes were coated with LPS and D₉ inhibitor then reacted with guinea pig serum (absorbed

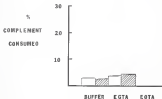
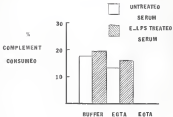
TABLE 1

Quantities of Virus Adsorbed to Cells in a 10-min. Incubation
 of Virus Adsorbed GP Serum
 to Uninfected and LPS-Coated Sheep Erythrocytes

Virus Quantities	Percent Adsorbed	
	E-absorbed serum	E-LPS-absorbed serum
0	—	—
10 ⁶ LPS absorbed	25.3	16.89
10 ⁷ LPS absorbed	42.9	6.35

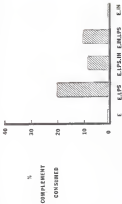
Figure 8

Comparison of compliance in normal and E-LPS treated groups plus serum by three phase nitrogenella labelled LPS detected by the triple and sequential procedures both, 100%, and saline treated sera were treated with the LPS preparation one hour at 37°C, then sedimented after which the C800 filters were determined in the nitrogen and carbon monoxidized sera. The top figure represents the profile of the normal treated sera and the lower figure, the biopsied treated sera



cells [32] (control) [33] (no fluid) all cells were lysed. The third group was coated with LPS only (L-LPS), a second group was coated with (DL-LPS) first, then was treated with the IL-1 inhibitor (IL-LPS-IL1), a third group was treated with IL-1 inhibitor first, followed by the LPS-receptor (IL-LPS-IL1), a fourth group was treated with IL-1 inhibitor only (IL-IL1), and, a fifth group of cells [34] was treated with PBS under the same conditions and served as the control. The efficiency of the LPS coating procedure in the presence and absence of the IL-1 inhibitor was evaluated by assaying the five groups of cells and their respective supernatant fluids for LPS activity, employing the hemagglutination assay (described in Roberts and Pettibone) as determined with antisera to *Salmonella typhimurium* group D. The results of these assays indicated that all of the LPS treated cells adsorbed equal quantities of LPS in the presence and absence of the IL-1 inhibitor. The complement consumption profiles of the five groups of erythrocytes treated with D-LPS (control) appeared similar (Fig. 1) and are presented in Figure 1C. It can be seen that the presence of IL-1 inhibitor on the cell caused about 50% reduction in the LPS mediated hemolytic action of complement. Of particular interest was the fate of the cells when complement was treated with DL + LPS coated erythrocytes, right blood/tot cells were lysed. This was in contrast to the case where complement was mixed with D-LPS and the cells were completely lysed. These results suggested the possibility that complement activation had taken place, but that cells were protected from lysis by the presence of IL-1 inhibitor on the membrane.

Depletion of complement in *E-LPI* sensitized guinea pigs was by *LPI* and/or *D* inhibitor coated sheep erythrocytes. One batch of sensitizing 10^8 sheep erythrocytes (*E*) either untreated or treated with *LPI* (*E-LPI*), *D* inhibitor (*I-I*) or *LPI* and *D* inhibitor (*D-LPI-I*) were incubated with 3-5 ml of guinea pig serum at 37°C for 1 hr. Residual complement hemolytic activities were assayed and the % of the available complement consumed was calculated.



DISCUSSION

The experiments reported here have demonstrated that extracts from human erythrocyte membranes possessing LPI-receptor activity determined by the method of Springer et al. (15) were also capable of complement mediated lysis. The anticomplementary activity of these extracts was demonstrated to share many of the properties of the IR inhibitor previously described by Hoffmann (16).

Data are presented which strongly suggest that the two biological activities are closely associated, but separable. Evidence for this was provided by the results of five different experimental approaches in the analysis of Springer human erythrocyte stromal extracts. The first was based on the chromatographic properties on Sepharose 4B and DEAE-Sephadex where slight differences between the elution patterns and recoveries of the two activities were observed. The second piece of evidence came from the electrophoretic profile of the crude and partially purified extracts on 7.5% polyacrylamide gels under non-reducing conditions. The IR inhibitor activity was shown to cover a fairly large area at the top third of the gel with the LPI-receptor activity being localized in a narrow, single band with a peak of activity near the top of the gel. A third line of evidence was based on the redistribution and separation of the two activities into different phases when the pH during the crude stromal (stromal) extraction procedure

shifted from 5.2 to 5.3. The fourth approach, based on the high affinity of the Ia inhibitor for the membranes of sheep erythrocytes, demonstrated that the Ia inhibitor activity was partially removed leaving the LPS-receptor activity unchanged by the treatment of the stromal extracts with sheep erythrocytes. Finally, evidence was presented indicating that the binding specificity of sheep erythrocytes for LPS of gram negative bacteria is localized in a lipid moiety of the crude stromal extracts and is free of all detectable Ia inhibitor activity.

It should be emphasized, however, that these experiments cannot exclude the possibility that both activities may be associated to the same macromolecule with the differences reported here being a consequence of experimental manipulation. That the two activities may be a function of a single macromolecule is certainly a major possibility. Springer et al. (14) in assessing the chemical and physical properties of a homogeneous preparation of the LPS-receptor, observed that both citraconylation and dissociating polyacryamide gel electrophoresis under standard conditions yielded two fragments, one of which absorbed significantly only at 330 nm. Backconjugation of the citraconylated fragment restored high LPS-receptor activity to half one of the fragments. These studies are only suggestive and do not permit a decision as whether the activities are on the same molecule however.

In contrast, the data obtained from sheep erythrocytes which completely lack Ia inhibitor, but which possess LPS-receptor activity, would support the finding that the two macromolecules may be distinctly different. However, the evidence could support that the LPS-receptors

on sheep cells differ from those observed on human cells, since they are confined to the T and M activity.

Good purification of the LPS-receptor and its inhibitor activities following DEAE-cellulose ion-exchange chromatography is indicated by the quantitative data tables. These results would suggest, however, that as a preparative purification step it should be modified to encourage higher yields of the two activities.

The observation that sheep erythrocyte membranes possess interferes with receptor specificity for LPS is not surprising for it has long been known that sheep cells could be agglutinated by the presence of LPS of gram negative bacteria and that these agglutinated cells are readily lysed in the presence of hemolytic antibodies to LPS and complement (40). In contrast, hemolysis was not observed when LPS treated human erythrocytes were treated under the same conditions. This raises the possibility that human erythrocytes, a natural source of the M inhibitor even when treated with LPS, are extremely resistant to LPS mediated lysis brought about by the presence of inhibitor molecules.

The findings reported here generally agree with those of Phillips et al. (41) indicating that LPS treated sheep erythrocytes can activate the complement system in the presence of natural antibodies to the LPS. In contrast to their results, however, erythrocytes coated with a preparation of LPS extracted by the procedure of Weinstein were shown to be capable of activating the complement system in the absence of natural antibodies to LPS. Additionally, fluid phase LPS extracted by the Levine method (LPS-bound) was shown to be a more effective activator of the

complement system than LPS extracted by the heat-kill procedure (LPS-*heat-kill*) in the presence and absence of natural antibodies to LPS. However, since LPS extracted by the Boivin became cell associated, its capacity to activate the complement system in the absence of natural antibodies is greatly diminished. This is significant because LPS activation of the complement system by an antibody-independent mechanism requires either an exposed lipid A moiety or polysaccharide side chain (HUB). This would support the fact that the orientation of LPS-moieties on the membrane may be different from that of LPS-*heat-kill*, resulting in the masking of active sites necessary for the activation of complement.

The fact that different preparations of LPS from the same species, when coated onto the surface of sheep erythrocytes, activated the complement system to different degrees and by different pathways, depending on the presence or absence of natural antibodies to LPS, introduces the possibility that the LPS activation of complement may require substances other than the LPS molecule alone. This especially appears to be true since LPS extracted by the heat-kill procedure, which was shown to activate complement in the presence and absence of natural antibody to LPS, is known to contain less proteins and lipoproteins than LPS extracted by the Boivin procedure.

Having the IR inhibitor on LPS-treated sheep erythrocytes reduced the ability of erythrocytes to consume complement. The fact that the cell is protected even when complement is activated suggests that the IR inhibitor may occupy specific sites on the red cell membrane rendering it resistant to osmotic lysis, but also leaving it available

to partially inhibit tissue lysis. This may be due either to the masking of LPS-receptors resulting in less LPS uptake (demonstrated not to be the case here), or the presence of CR proteins on the cell membrane thus blocking the initiation of the membrane attack system of serum complement. An analogous site in human erythrocytes is already occupied by the Ii inhibitor rendering this cell naturally immune to LPS mediated lysis.

The findings presented here have led us to hypothesize that a necessary criterion for resistance to LPS induced complement mediated lysis would be the localization of LPS-receptor and Ii inhibitor molecules on the same membrane. This would imply that any red cell devoid of Ii inhibitor molecules would be far more susceptible to the cytolytic action of LPS activated complement.

LPS of gram negative bacteria are potent activators of the complement system. It is of great clinical interest, therefore, that substances on the surface of erythrocytes which bind LPS are found directly associated with materials capable of inhibiting complement mediated lysis. The symptoms of several infectious diseases, such as typhoid fever, have been observed to include very intensive erythrophagocytic activity by macrophages of lymph nodes. The consequences of this observation could have great clinical importance. Erythrocytes coated with LPS, in contact with serum complement, would naturally lead to activation of the complement system followed by increased phagocytosis with eventual cytolysis. This would lead naturally to an amplification of the activation of the complement cascade resulting in either clearance or a heightened inflammatory response.

REFERENCES

1. Barten, A. 1968. The antigenic sensitization of avian erythrocytes with *Salmonella gallinarum* polysaccharide. *Immunology* **2**:308.
2. Leite, L., V. K. Shewlin and R. L. Nussenzweig. 1968. Physical, chemical, and immunological properties of LPS released from *Escherichia coli* by ethylene diamine tetraacetate. *J. Biol. Chem.* **243**:4364.
3. Weil, R. J. and L. Leite. 1970. Release of LPS by *Escherichia coli* resistant to the permeability increase induced by ethylenediamine tetraacetate. *J. Biol. Chem.* **245**:1588.
4. Kaper, J. B. C. and R. B. Kellner. 1971. Bacterial Toxins. Wolstein, G., S. Kaper, and L. J. Apt, eds. Academic Press, New York. 361-4, pp. 291-321.
5. Harrison, G. C. and L. Leite. 1972. Fractions of lipopolysaccharide from *Escherichia coli* O157 prepared by two extraction procedures. *J. Biol. Chem.* **247**:2911.
6. Gidycz, R. J. 1969. Structure and biosynthesis of the bacterial ap⁺ cell. *Ann. Rev. Biochem.* **38**:505.
7. Luderitz, O., V. L. Galanos, K. Nowinski, E. T. Rietschel, G. Rosenfelder, H. Stone, and O. Hartshel. 1973. 'Lipid A.' Chemical structure and biological activity. *J. Infect. Dis.* **133**:817.
8. Harrison, G. C. and L. F. Wise. 1972. Activation of the classical and properdin pathways of complement by bacterial lipopolysaccharides (LPS). *J. Immunol.* **118**:342.
9. Springer, George F. and J. E. Abo. 1973. Endotoxin-binding substances from human leukocytes and platelets. *Infect. and Imm.* **12**:479.
10. Sawyer, J., A. Sawyer, and L. Timmons. 1975. Endotoxin sensitive membrane components of human platelets. *Science* **228**:129.
11. Kane, R. A., J. E. May, and R. B. Fress. 1972. Interaction

of the alternative and alternate complement pathway with endotoxin lipopolysaccharide - effects on platelets and blood coagulation. *J Clin Invest* 52:179

12. Holt, C. J. and D. B. Davis. 1964. Relationships between susceptibility of mice to heat-killed *Salmonella* and endotoxin and the size affinity of their red blood cells for killed organisms. In R. Landy and S. Aron, eds. *Bacterial Endotoxins*. Institute of Microbiology, Rutgers, the State University, New Brunswick, N.J. 440-447.
13. Springer, G. F., C. T. Wang, J. B. Nichols, J. B. Sheer. 1968. Relations between bacterial lipopolysaccharide structures and those of human cells. *Am. J. Hyg.* 91: 131-144.
14. Springer, G. F., A. Shender, E. A. Rapoport, and C. Suter. 1970. Specific inhibition of endotoxin coating of red cells by a human erythrocyte membrane component. *Infect Immun* 1: 35.
15. Springer, G. F., J. C. Adge, A. Berkmanovitz, and J. B. Ruthe. 1972. Functional aspects and nature of the lipopolysaccharide receptor of human erythrocytes. *J. Infect. Dis.* 218: 850.
16. Springer, G. F., J. C. Adge, A. Berkmanovitz, and B. Jirymason. 1974. Properties and activity of the lipopolysaccharide receptor from human erythrocytes. *Science* 182: 1476.
17. Liss, R., B. Eftar Sorensen, and B. Starck. 1971. Interaction of the first (C1) and second (C2) and fourth (C4) component of complement with different preparations of bacterial lipopolysaccharide and with lipids. *J. Immunol.* 116:815.
18. Manning, N. B., J. C. Hansen, R. E. Ishler, and J. B. Palmer. 1969. Distribution and clearance of circulating endotoxins. *J. Clin. Invest.* 52: 71.
19. Morgenstern, L. E., B. Seyferman, and J. C. Phillips. 1973. Activation of complement by endotoxin. *J. Infect. Dis.* 232: 548.
20. Møller-Gerhard, H. J. 1974. Complement. *Am. Rev. Biochem.* 52: 617.
21. Sirtes, B. and H. J. Møller-Gerhard. 1976. The alternative pathway of complement activation. *Advances in Immunol.* 24: 1-100.
22. McConnell, J. and P. J. Lachman. 1971. Complement receptors and cell associated complement components. *Immunol. Rev.* 6: 115.

23. Rifkinen, L., M. D. Schneider, L. Blum, and L. Burg. 1963. Properties system and instability. VI. Interaction of the properdin system with polysaccharides. *Science* 122: 889.
24. Maudel, L. B., E. Schneider, and P. H. Napp. 1964. Self-complementary action of endotoxin. *Proc. Soc. Exp. Biol. Med.* 117: 818.
25. Starick, M. P., G. Ritter-Sommert, A. Korte, B. Haddara, C. Salmons, and E. J. Hietzerfel. 1973. Analysis of bypass activation of C3 by endotoxin LPS and loss of this potency immunology. 28: 121.
26. Marcus, R. I., R. S. Stein, and T. M. Mayer. 1971. An alternative complement pathway. C3 cleaving activity not due to C4, is on endotoxin lipopolysaccharide after treatment with guinea pig serum, relation to properdin. *Proc. Natl. Acad. Sci.* 68: 1391.
27. Salmons, C. C., T. Hietzerfel, E. Gubersitz, and G. Hietzerfel. 1970. Interaction of lipopolysaccharides and lipid A with complement. *Eur. J. Biochem.* 13: 743.
28. Heller, G. and S. Michael. 1973. Frequency of antigen-sensitive cells to thymus-independent antigens. *Cell. Immunol.* 7: 389.
29. Cooper, R. B. and D. C. Harrison. 1973. Binding and activation of the first component of human complement by the lipid A region of lipopolysaccharides. *J. of Immunol.* 120: 1380.
30. Stein, G. and R. J. Heller-Barbard. 1971. The C3 activator system. An alternative pathway of complement activation. *J. Exp. Med.* 134: 905.
31. Ley, H. R. and E. Rausenberger. 1965. Receptors for complement on leukocytes. *J. Exp. Med.* 122: 941.
32. Snyderman, R. and H. C. Pike. 1973. Interaction of complex polysaccharides with the complement system. Effect of calcium depletion on terminal complement activation. *Infekt. Immun.* 11: 875.
33. Pargan, D., T. B. F. Austin, and S. Ruddy. 1973. Formation of a hemolytically active cellular intermediate by the interaction between properdin factors B and D and the activated third component of complement. *J. Exp. Med.* 138: 1595.

3. Paros, R., J. L. Jaffe, and R. M. Fayer. 1972. An alternative complement pathway: C3 cleaving activity not due to C3, C5, or C6. In *Immunology: Tissue Transplantation and Immunology*, with essays and topics, relation to properties (complement components). Proc. Natl. Acad. Sci. USA. 69:1211.
4. Fayer, R. M., R. M. Paros, R. C. Colley, J. S. Berges, and R. M. De Fayer. 1972. C3 convertase in human serum isolated with 197S. *J. Immunol.* 109:807.
5. Fayer, R. M. 1972. Mechanism of cytotoxicity by complement. *Proc. Natl. Acad. Sci. USA.* 69:2964.
6. Jansen, J. 1947. Anaphylatoxins and its relation to the complement system. *Science* 100:1123.
7. Irie, H., R. Snyderman, E. Friedman, A. Mallory and R. Fayer. 1968. Characteristic and anaphylatoxin fragments released from the fifth component of guinea pig complement. *Science*, 162:867.
8. May, J. E., R. A. Saxe, and R. M. Fayer. 1972. Immune adherence by the alternative complement pathway. *Proc. Soc. Exp. Biol. Med.* 141:287.
9. Kasper, L. 1972. Leukocyte sensitizing factor: A non-lysozymal activity derived from the third component of complement. *Int. J. Immunol.* 2:360.
10. May, J. E. and R. M. Fayer. 1972. Complement mediated tissue damage: Contribution of the classical and alternate complement pathways in the Porcine reaction. *J. Immunol.* 109:1217.
11. McTier-Cleburne, R. J. and J. A. Lopez. 1965. C7 esterase effect on activity and physicochemical properties of the fourth component of complement. *J. Exp. Med.* 121:879.
12. Ruddy, S. and C. F. Austen. 1975. C5b inactivator of man. II. Properties of anti serum of fluid phase C5b. *J. Immunol.* 105:792.
13. Reiser, E. 1966. Bacterial hemagglutination and hemolysis. *Bacteriol. Rev.* 30:164.
14. Weisbach, O. and J. Kain. 1965. Bacterial lipopolysaccharides: Extraction with alcohol and further purification of the products. In *Methods in Carbohydrate Chemistry*, R. L. Whistler, ed. Academic Press, New York. Vol. 5, p40.

46. Pollinger, J. M., R. Seydman, and S. C. Bergsbuhen. 1977. Activation of complement by antibodies: A role for C2, C3, C4 and C5 in the consumption of terminal complement components by antibody-coated erythrocytes. *J. of Immunol.* 128: 334.
47. Pap, J. E., R. A. Kane, and R. M. Frank. 1970. Host defense against bacterial endotoxemia - Contribution of the early and late components of complement to detoxification. *J. Immunol.* 109: 883.
48. Gewurz, H., R. S. Stein, and S. C. Bergsbuhen. 1968. Interactions of the complement system with endotoxic *Triglopor-saccharum*. Consumption of each of the six terminal complement components. *J. Exp. Med.* 128: 1049.
49. Pap, J. E., R. A. Kane, and R. M. Frank. 1972. Immune adherence by the alternate complement pathway. *Proc. Soc. Exp. Biol.-Med.* 139: 337.
50. Hoffmann, Edward H. 1968. Inhibition of complement by a substance isolated from human erythrocytes - extraction from human erythrocyte ghosts. *Immunochim.* 3: 231.
51. Hoffmann, E. H., K. L. Cheek, L. J. Tonks, and C. H. Rank. 1964. Resistance of sheep erythrocytes to immune lysis by treatment of the cells with a human erythrocyte extract. Studies on the site of inhibition. *J. of Immunol.* 122: 1627.
52. Hoffmann, E. H. 1963. Inhibition of complement by a substance isolated from human erythrocytes. II. Studies on the site and mechanism of action. *Immunochim.* 8: 450.
53. Hoffmann, E. H. and R. M. Cillipier. 1973. Extraction of complement inhibitory factors from the erythrocytes of non-human species. *J. of Immunol.* 121: 546.
54. Mason, H. L. and J. H. Sachs. 1968. Immune lysis of normal and Hb red cells. I. Sensitivity of Hb cells. *J. Clin. Invest.* 45: 236.
55. Sakai, E. A. and H. H. Meyer. 1967. Complement and complement proteins. In Immunological Investigations, Charles C. Thomas, Springfield, Ill. p145.
56. Springer, G., E. F. Rausl, and H. Tschreyer. 1969. Isolation and properties of human blood-group M and manganin-Vp antigens. *Biochem.* 8: 3234.

87. Fauser, R. B. 1981 Site Electrophoresis and Related Techniques of Polyacrylamide Gel Electrophoresis. 2nd revised ed. GUSTO de Gruyter, Berlin, 87.
88. Rotim, R. J. 1968 Extraction of bacterial O-antigen (endotoxin). Acas, Daresbury, Immunobiology. 4. Macmillan, ed. Springer-Verlag, New York. 1968. 525.
89. Wilson, R. A., J. Jensen, J. Sjoh, and E. Timmer. 1968. Methods for the separation, purification, and measurement of nine components of hemolytic complement in guinea pig serum. Immunology. 11.
90. Ruddy, S. and R. F. Austen. 1967. A microhemetic assay for the fourth component of complement in whole human serum using EACD and functionally pure human serum complement. J. Immunol. 100:182.
91. Ruddy, S. and R. F. Austen. 1968. C3 inactivator of man. I. Hemolytic measured by the inhibition of anti sheep E1. J. Immunol. 100:583.
92. Hesse, E. and H. J. Lipp. 1967. Immune hemolysis. A simplified method for preparation of EACs with guinea pig or with human complement. J. Immunol. 100:665.

Continued

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I certify that I have read this study and that in my opinion it conforms to acceptable standards of scholarly presentation and is fully adequate, in scope and quality, as a dissertation for the degree of Doctor of Philosophy.


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I hereby state, I do hereby certify, that in my opinion it conforms to acceptable standards of scholarly presentation and is fully adequate, in scope and quality, to be accepted as a dissertation for the degree of Doctor of Philosophy.


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I certify that I have read this study and that in my opinion it conforms to acceptable standards of scholarly presentation and is fully adequate, in scope and quality, to be accepted as a dissertation for the degree of Doctor of Philosophy.


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